Introduction

Most computers around nowadays have some sort of molecular modeling/viewing program available in the public domain or commercially. I needed such a beast for my research (molecular biophysics), and couldn't find one for the NeXT ... hence MolViewer was born.

This is the second release of MolViewer. As promised, complete source is included this time. There are quite a few new features in this release, but I haven't put in everything I'd hoped to. If I have time I'll put out another release in the next month or so ... The bug in QRM is still a problem for large molecules, but the workaround still seems to work fairly well.

Again, I would greatly appreciate suggestions and bug reports, now that I've got my Masters I have a lot more spare time to work on this. Before making suggestions or bug reports, please read the history file (in help) for the list of known bugs and future plans. Thank you, and enjoy ...

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